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Reviewer: Durreshwar Anjum

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Application No: 10565591 Version No: 1.0

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No. of SeqIDs Defined: 5
Actual SeqID Count: 5

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SEQUENCE LISTING

<110> KARAOLIS, David K.R.

<120> METHOD FOR ATTENUATING VIRULENCE OF MICROBIAL PATHOGENS AND FOR
INHIBITING MICROBIAL BIOFILM FORMATION

<130> KARAOLIS1 PCT

<140> 10565591

<141> 2007-05-25

<150> NOT YET ASSIGNED

<151> 2004-07-22

<150> 60/490,029

<151> 2003-07-25

<160> 5

<170> PatentIn version 3.2

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Thr Ser His Ser Pro Phe Phe Phe Ala Ile Leu Asn Asp Gln His Gln	
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Glu Glu Met Val Gly Met Ser Asp Ser Gln Val Leu Gly Glu His Phe	
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tat cgc cat ctc aaa ccg ttt tac gaa cgt gcg ttt aac aac gag cat	240
Tyr Arg His Leu Lys Pro Phe Tyr Glu Arg Ala Phe Asn Asn Glu His	
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att gag tcc gag ctg acc ctc agc gaa atc gac ctc gaa acc agc tta	288
Ile Glu Ser Glu Leu Thr Leu Ser Glu Ile Asp Leu Glu Thr Ser Leu	
85 90 95	

cac ttt tct ctc tcc ccc atc atg atc aac gat cgg gtg caa tac ctt	336
His Phe Ser Leu Ser Pro Ile Met Ile Asn Asp Arg Val Gln Tyr Leu	
100 105 110	
gta ttc cac gcg att gat acc tca gaa aag cag att tta gtg cgc tct	384
Val Phe His Ala Ile Asp Thr Ser Glu Lys Gln Ile Leu Val Arg Ser	
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Leu Glu Glu Ser Glu Ser Lys Tyr Ala Leu Leu Thr Thr Leu Leu Pro	
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Ser Ala Ala Arg Leu Leu Gly Phe Asp Asp Ala Gln Lys Leu Leu Gly	
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Glu Asn Leu Ser Arg Leu Phe Ile Asp Glu Lys Thr Lys Thr Val Phe	
180 185 190	
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Ser Ser Gln Leu Ala Ser Leu Leu Thr Glu Lys Pro Leu Val Cys Leu	
195 200 205	
acc ggg cca agg tgt ggg ttt gaa cgg aaa atc cag tta cac gca ggt	672
Thr Gly Pro Arg Cys Gly Phe Glu Arg Lys Ile Gln Leu His Ala Gly	
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tgc acc tct tta ctc ggt aat cag tcg cag ttg atc tta ttg caa gat	720
Cys Thr Ser Leu Leu Gly Asn Gln Ser Gln Leu Ile Leu Leu Gln Asp	
225 230 235 240	
gcc gat gaa gcc cca aaa cag ttt tct gcg acc act caa gtc gat gcg	768
Ala Asp Glu Ala Pro Lys Gln Phe Ser Ala Thr Thr Gln Val Asp Ala	
245 250 255	
cat att gat agc ctc act ggg ctg tat aac cga cac ggg ttt acc aag	816
His Ile Asp Ser Leu Thr Gly Leu Tyr Asn Arg His Gly Phe Thr Lys	
260 265 270	
cgc tta gag cag tgc atc caa aat gag acg cct ttg gtt atg ctc tat	864
Arg Leu Glu Gln Cys Ile Gln Asn Glu Thr Pro Leu Val Met Leu Tyr	
275 280 285	
ctg gac att gat aac ttc aaa aac atc aat gac tct ctc ggc cat cac	912
Leu Asp Ile Asp Asn Phe Lys Asn Ile Asn Asp Ser Leu Gly His His	
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Ile Gly Asp Lys Val Ile Lys Glu Val Ala Ala Arg Leu Lys Arg Leu	
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Leu Pro Gln Gln Ala Val Leu Gly His Leu Gly Gly Asp Glu Phe Gly	
325 330 335	
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Leu Ile Leu Pro Glu Pro Glu His Asn Arg Ser Ala Glu Met Leu Ala	
340 345 350	
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Asp Arg Ile Ile Ser Leu Ile Asn Gln Pro Phe Asp Leu His His Phe	
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agt aag cgt tta gct tgt tcg att ggc agc gtg cgt tat ccc ggt gac	1152
Ser Lys Arg Leu Ala Cys Ser Ile Gly Ser Val Arg Tyr Pro Gly Asp	
370 375 380	
ggc aat gat gct cgc gta tta ctg caa aat gcc gat acc gcg atg tat	1200
Gly Asn Asp Ala Arg Val Leu Leu Gln Asn Ala Asp Thr Ala Met Tyr	
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Met Asn Lys Glu Ala Arg Met Arg Leu Trp Leu Glu Ile Glu Leu Gln	
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aaa gcg cta caa caa aac ggc cta gaa gtg tgg tac caa ccg aaa gtc	1344
Lys Ala Leu Gln Gln Asn Gly Leu Glu Val Trp Tyr Gln Pro Lys Val	
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aac gcg cgt gat ttt agc atc aat ggc gca gaa gcc ttg gta cgc tgg	1392
Asn Ala Arg Asp Phe Ser Ile Asn Gly Ala Glu Ala Leu Val Arg Trp	
450 455 460	
aaa cat ccc gtt gaa ggc tat atc agc cca ggt gct ttc att ccc gtt	1440
Lys His Pro Val Glu Gly Tyr Ile Ser Pro Gly Ala Phe Ile Pro Val	
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Ala Glu Lys Ala Gly Leu Ile Glu His Leu Gly Arg Val Val Met Arg	
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gaa gtc ttc gcg acc gtc aag cgc tgg aag cta caa ggc att tta ccc	1536
Glu Val Phe Ala Thr Val Lys Arg Trp Lys Leu Gln Gly Ile Leu Pro	
500 505 510	
gga cgt gtg gcg atc aac atc tcc ccc gag cag ttt ggc aat cct caa	1584
Gly Arg Val Ala Ile Asn Ile Ser Pro Glu Gln Phe Gly Asn Pro Gln	
515 520 525	
ctg att gat tat tta gaa aaa cta ctg cga aca act ggg cta gat ccc	1632
Leu Ile Asp Tyr Leu Glu Lys Leu Leu Arg Thr Thr Gly Leu Asp Pro	
530 535 540	
aac aac atc aca ttt gaa ctg acc gaa agt gtg gtg atg agc gat agt	1680
Asn Asn Ile Thr Phe Glu Leu Thr Glu Ser Val Val Met Ser Asp Ser	

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Glu His Thr Gln Gln Met Leu Asn Ala Ile Lys Lys Leu Gly Phe Thr				
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ttg tca att gat gac ttc ggt aca ggt tac tcg tcg ctg gct tat tta				1776
Leu Ser Ile Asp Asp Phe Gly Thr Gly Tyr Ser Ser Leu Ala Tyr Leu				
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gct cgc ttc ccg atc gat gag ctc aaa atc gac cgc gcg ttt atc agt				1824
Ala Arg Phe Pro Ile Asp Glu Leu Lys Ile Asp Arg Ala Phe Ile Ser				
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aat atc gac act cta ccc aaa cag ctc acg gtg att gaa aac atc att				1872
Asn Ile Asp Thr Leu Pro Lys Gln Leu Thr Val Ile Glu Asn Ile Ile				
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aat ttg ggg cgc tca ctg aac ctg acc gta gtt gca gaa gga gta gaa				1920
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act cag caa caa gcc act tta ctc tcc aac cta aat tgc cac tcc atc				1968
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caa ggc ttc cat ttt tat cgc cca caa ccg aag cac gaa gtg gaa gag				2016
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Glu Glu Met Val Gly Met Ser Asp Ser Gln Val Leu Gly Glu His Phe
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Ile	Glu	Ser	Glu	Leu	Thr	Leu	Ser	Glu	Ile	Asp	Leu	Glu	Thr	Ser	Leu	85	90	95	
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Asp	Gly	Leu	Met	Met	Val	Glu	Asn	Asp	Cys	Ile	Ile	Ser	Ala	Asn	Pro	145	150	155	160
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Glu	Asn	Leu	Ser	Arg	Leu	Phe	Ile	Asp	Glu	Lys	Thr	Lys	Thr	Val	Phe	180	185	190	
Ser	Ser	Gln	Leu	Ala	Ser	Leu	Leu	Thr	Glu	Lys	Pro	Leu	Val	Cys	Leu	195	200	205	
Thr	Gly	Pro	Arg	Cys	Gly	Phe	Glu	Arg	Lys	Ile	Gln	Leu	His	Ala	Gly	210	215	220	
Cys	Thr	Ser	Leu	Leu	Gly	Asn	Gln	Ser	Gln	Leu	Ile	Leu	Leu	Gln	Asp	225	230	235	240
Ala	Asp	Glu	Ala	Pro	Lys	Gln	Phe	Ser	Ala	Thr	Thr	Gln	Val	Asp	Ala	245	250	255	
His	Ile	Asp	Ser	Leu	Thr	Gly	Leu	Tyr	Asn	Arg	His	Gly	Phe	Thr	Lys	260	265	270	
Arg	Leu	Glu	Gln	Cys	Ile	Gln	Asn	Glu	Thr	Pro	Leu	Val	Met	Leu	Tyr	275	280	285	

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Ile Gly Asp Lys Val Ile Lys Glu Val Ala Ala Arg Leu Lys Arg Leu
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Leu Pro Gln Gln Ala Val Leu Gly His Leu Gly Gly Asp Glu Phe Gly
325 330 335

Leu Ile Leu Pro Glu Pro Glu His Asn Arg Ser Ala Glu Met Leu Ala
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Asp Arg Ile Ile Ser Leu Ile Asn Gln Pro Phe Asp Leu His His Phe
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Ser Lys Arg Leu Ala Cys Ser Ile Gly Ser Val Arg Tyr Pro Gly Asp
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Gly Asn Asp Ala Arg Val Leu Leu Gln Asn Ala Asp Thr Ala Met Tyr
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Glu Ala Lys Glu Arg Gly Arg Asn Arg Leu Ile Lys Phe Asn Asp Gln
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Met Asn Lys Glu Ala Arg Met Arg Leu Trp Leu Glu Ile Glu Leu Gln
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Lys Ala Leu Gln Gln Asn Gly Leu Glu Val Trp Tyr Gln Pro Lys Val
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Asn Ala Arg Asp Phe Ser Ile Asn Gly Ala Glu Ala Leu Val Arg Trp
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Lys His Pro Val Glu Gly Tyr Ile Ser Pro Gly Ala Phe Ile Pro Val
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515

520

525

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Asn Ile Asp Thr Leu Pro Lys Gln Leu Thr Val Ile Glu Asn Ile Ile
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